

Phylogenetic Analysis of Bird-Virulent West Nile Virus Strain, Greece

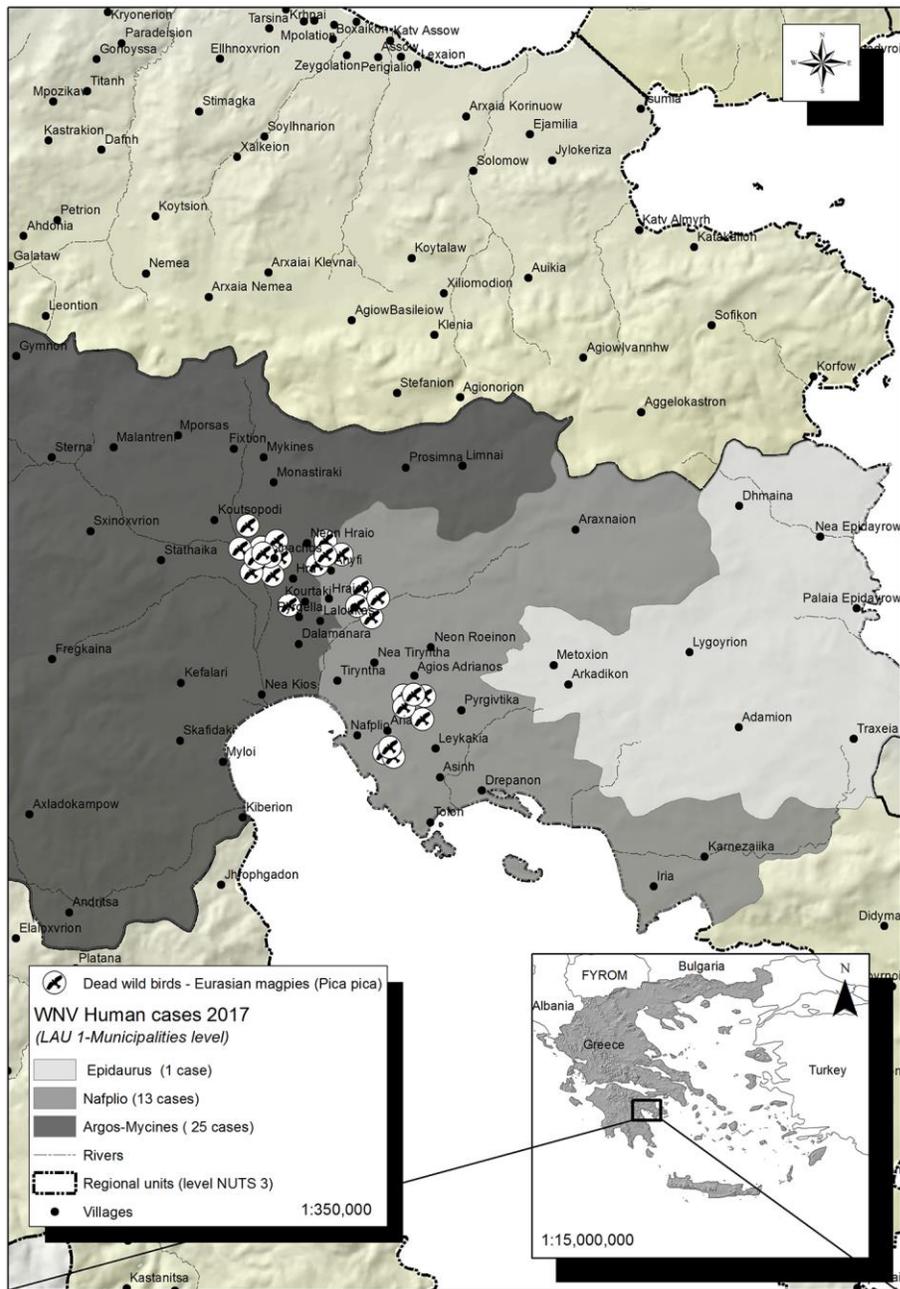
Appendix

Appendix Table. Primers used in analysis of bird-virulent strain of West Nile virus, Greece

| Primer name | Sequence primer (5' →3') | Nea Santa-Greece-2010 nt position | Bibliography |
|-------------|--------------------------|-----------------------------------|--------------|
| F1 | AGTAGTTCGCCTGTGTGAG | 1–19 | This study |
| R1 | GCCTTCCAGTACCAGATCAA | 1028–1047 | |
| F2 | TDGGAATGAGYAACAGRGAC | 979–996 | (1) |
| R2 | TGRTTDATECTGBTGWTCTCCT | 2130–2150 | |
| F3 | GTTCTCAAGCAATACTGTGAAG | 1785–1806 | (2) |
| R3 | TTCCACCACGACACTCAA | 2725–2742 | |
| F4 | GGAAGCCATTAAGGATGAG | 2673–2691 | (2) |
| R4 | AAGAACACGACCAGAAGG | 3561–3578 | |
| F5 | AGAATGGCTGTTGGTATGG | 3446–3464 | (2) |
| R5 | CGTCCTCTCAATCCACAT | 4372–4389 | |
| F6 | GCAGAACTTGACATAGACTC | 4282–4301 | (2) |
| R6 | TACAGCCGTCCTCAATCT | 5257–5274 | |
| F7 | ACCTGAAATGTTGAGGAAGA | 5151–5170 | (2) |
| R7 | CGGCATGTTGATGTTGTC | 6100–6117 | |
| F8 | CATCACCACAGACATATCAGA | 5829–5849 | (2) |
| R8 | AGCCATCCAGCAGAAGAA | 6742–6759 | |
| F9 | CGGACGCCCTTCAGACAATA | 6611–6630 | (2) |
| R9 | TCGTTACGGCATAGAGCGAC | 7044–7059 | |
| F10 | CCTGTTGGGACACAAGCCA | 6960–6978 | This study |
| R10 | CCAGATACAGAAAAGAAGCCATCA | 7757–7780 | |
| F11 | CAAGAGGGGTGGAGCCAAG | 7677–7695 | This study |
| R11 | AGGAGAGGCTTCCCTACTGC | 8479–8498 | |
| F12 | GTGAACATGACAAGCCAGGTG | 8380–8400 | This study |
| R12 | CTGAGTTCTTCTACCCAGCC | 9179–9199 | |
| F13 | GAGTTCGGCAAAGCTAAAGGC | 9079–9099 | This study |
| R13 | ACACTCTGTTCCAGACTGCG | 10128–10147 | |
| F14 | TGGATGACGACGGAAGACATG | 10105–10125 | (3) |
| R14 | GGGTCTCTTAACCTCTAGT | 10825–10845 | |

References

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Appendix Figure. Area in which dead Eurasian magpies were collected during July and August 2017 in a study of bird-virulent West Nile virus, Greece.